

Holding Chandra  
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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

ATTY.'S DOCKET: RUBINSTEIN=7

In re Application of:	)	Conf. No.: 2828
	)	
Menachem RUBINSTEIN et al	)	Art Unit: 1646
	)	
Appln. No.: 10/070,295	)	Examiner: G. Chandra
	)	
Date Filed: March 5, 2002	)	Washington, D.C.
	)	
For: USE OF LEPTIN IN	)	
INHIBITION OF ENDOTHELIAL	)	
CELL PROLIFERATION	)	

DECLARATION UNDER 37 CFR §1.132

Honorable Commissioner for Patents  
U.S. Patent and Trademark Office  
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Randolph Building, Mail Stop  
401 Dulany Street  
Alexandria, VA 22314

Sir:

I, Menachem RUBINSTEIN, declare and state as follows:

I am a co-inventor in the above-identified application and my educational and professional experience is presented in the curriculum vitae attached hereto.

I understand that the examiner has made an enablement rejection and has stated that, while the specification teaches leptin inhibiting angiogenesis in ob/ob mice, there is no guidance or any other example that leptin would inhibit angiogenesis in a normal mammal. Thus, the examiner takes the position that one of skill in the art is

not enabled for inhibiting angiogenesis in mammals that are normal and not ob/ob.

The examiner's logic in the lack of enablement rejection however is faulty. One of the most convincing ways to prove the function of any given protein is to generate a knockout mouse, which is unable to express this protein. It is then customary to administer this protein in order to show that its biological activity is restored. There is a vast body of scientific literature, which is based on this paradigm and it is an accepted method by virtually all life scientists. Although fa/fa rats and ob/ob mice are phenotypically analogous, it is well established that ob/ob mice can be cured by exogenous administration of leptin, whereas fa/fa rats do not respond to leptin whatsoever. Therefore, the two rodent models differ phenotypically in the most relevant issue in the present application, i.e., response to leptin.

The ob/ob strain is a spontaneously generated leptin knockout mouse. It has a single point mutation in the leptin gene, which disrupts its open reading frame, resulting in leptin deficiency. Otherwise, this strain is genetically perfectly normal. The experimental results presented in the instant application demonstrate that the addition of leptin to a leptin-deficient (ob/ob) mouse resulted in induction of angiopoietin 2 and inhibition of

angiogenesis in its adipose tissue. As discussed above, this proves the antiangiogenic function of leptin in adipose tissue (It should be noted that angiopoietin 2 acts differently in the presence or absence of VEGF, where in the absence of VEGF, it will induce vascular regression and endothelial cell death, as shown in Fig. 2 of Hanahan, *Science* 277(5322):48-50 (1997), a copy of which is attached). However, in the case of a normal mouse (not leptin-deficient), the administration of leptin to a normal mouse is like bringing coal to Newcastle. It would make a little or no difference since normal adipose tissue makes its own leptin.

The Cao et al. reference cited by the examiner demonstrates that leptin induces capillary growth in the cornea model, although less than VEGF (Fig. 1). In the adipose tissue, absence of leptin in ob/ob mice reduced the extent of fenestration as compared with wild type mice (Fig. 3). The extent of fenestration induced by VEGF in the cornea is greater than that induced by leptin (Fig 2), yet leptin induced more leakage (Fig.4). Fenestration and capillary growth are two distinct processes. Cao et al. however did not measure capillary growth in the adipose tissue, but rather explained their observation of fewer capillaries in the adipose tissue of ob/ob mice as an outcome of larger adipocyte size and not due to lack of leptin (see page 6932, second

column, first paragraph). Thus, Cao certainly does not teach or suggest that leptin induces capillary growth in adipose tissue, as asserted by the examiner.

In conclusion, one of skill in the art would indeed be enabled for inhibiting angiogenesis in mammals as presently claimed.

The undersigned declares further that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

Jan 22<sup>nd</sup>, 2008  
Date

Menachem Rubinstein  
Menachem RUBINSTEIN

## 1971CURRICULUM VITAE

Name : RUBINSTEIN, Menachem

Date & Place of Birth: 1945; Tel-Aviv, Israel.

Present home address: 1 Hohit St. Rehovot, Israel.

### EDUCATION:

- 1966 - 1968 B.Sc. in Chemistry and Biochemistry,  
The Hebrew University, Jerusalem, Israel.
- 1969 - 1970 M.Sc. in Biochemistry,  
The Hebrew University, Jerusalem, Israel.
- 1971 - 1975 Ph.D., Department of Organic Chemistry  
The Weizmann Institute of Science, Rehovot, Israel.  
Mentor: Prof. Abraham Patchornik.

### POSITIONS:

- 1976-1977 Postdoctoral Fellow, Dept. of Physiological Chemistry,  
Roche Institute of Molecular Biology.  
Nutley, New Jersey, USA.  
Mentor: The late Dr. Sidney Udenfriend
- 1978-1979 Visiting Scientist - Dept. of Physiological Chemistry and  
Pharmacology, Roche Institute of Molecular Biology.
- 1980-1983 Senior Scientist, Dept. of Virology,  
The Weizmann Institute of Science, Rehovot, Israel.
- 1983-1998 Associate Professor (with tenure) Dept. of Molecular  
Genetics,  
The Weizmann Institute of Science.
- 1987-1990 Chief Scientist (R&D manager), InterPharm Laboratories,  
Kiryat-Weizmann, Nes-Ziona, Israel.
- 1990-1991 Visiting Associate Professor, Dept. of Pharmacology  
New-York University, Medical Center, New York, U.S.A.
- 1991-2000 Head, Biological Services, The Weizmann Institute of Science.

- 1998- present      Professor, Dept. of Molecular Genetics,  
The Weizmann Institute of Science.
- 2002-2004      Head, The postdoctoral Fellowship program,  
The Feinberg Graduate School  
The Weizmann Institute of Science.
- 2006    2008      Chair  
The Scientific Council  
The Weizmann Institute of Science.

#### HONORS:

- 1975      Landau Prize for Ph.D. Thesis.
- 1980      Ionel Sanieel Career Development Chair.
- 1983      The Maurice and Edna Weiss Chair of Cytokine Research.
- 1996-1998      Member of the Editorial Board, Cytokines and Molec. Therapy.
- 1998-      Member of the Editorial Board, J. Interferon and Cytokine Res.
- 2000-      Member of the Editorial Board, Cytokine
- 2005      Recipient of the Milstein Award for IFN and Cytokine  
Research

#### PROFESSIONAL MILESTONES:

- |   |              |
|---|--------------|
| Development of HPLC methods for protein isolation and analysis            | 1977-1979    |
| Isolation of opioid peptides and their precursors                         | 1977-1978    |
| Isolation and characterization of human interferon alpha                  | 1978-1979    |
| Characterization of the human interferon- $\gamma$ receptor               | 1982-1986    |
| Identification of soluble IFN- $\gamma$ and IL-6 receptors in body fluids | 1988-1990    |
| Identification of a soluble LDL receptor as an antiviral protein          | 1989-1993    |
| Characterization and molecular cloning of the Type I IFN receptor         | 1992-1994    |
| Study of early events in signaling of the Type I IFN receptor             | 1995-1997    |
| Leptin receptor signaling and leptin as an angiogenic modulator           | 1996-2005    |
| Isolation and cloning of interleukin-18 binding protein                   | 1997-2003    |
| Regulation of IFN-gamma-induced gene expression                           | 2000-present |

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2. Rubinstein, M. and Patchornik, A. Polymers as chemical reagents. The use of poly(3,5-diethylstyrene) sulfonyl chloride for the synthesis of internucleotide bonds. *Tetr. Letters* 2881, 1972.
3. Hassner, A., Strand, G., Rubinstein, M. and Patchornik, A. Levulinic esters. An alcohol protecting group applicable to some nucleosides. *J. Amer. Chem. Soc.*, **97**, 1614, 1975.
4. Rubinstein, M., Amit, B. and Patchornik, A. The use of a light-sensitive phosphate protecting group for some mononucleotide syntheses. *Tetr. Letters* 1445, 1975.
5. Rubinstein, M. and Patchornik, A. Poly(3,5-diethylstyrene) sulfonyl chloride. A new reagent for internucleotide bond synthesis. *Tetrahedron* **31**, 1517, 1975.
6. Rubinstein, M. and Patchornik, A. A novel method for phosphodiester and internucleotide bond synthesis. *Tetrahedron* **31**, 2107, 1975.
7. Shechter, Y., Rubinstein, M., Becker, R. and Bohak, Z. Modulating of the enzymic activity of chicken pepsin by the covalent modification of its single SH group. *Europ. J. Biochem.* **58**, 123, 1975.
8. Rubinstein, M., Shechter, Y. and Patchornik, A. Covalent chromatography the isolation of tryptophanyl containing peptides by novel polymeric reagents. *Biochem. Biophys. Res. Comm.* **70**, 1257, 1976.
9. Shechter, Y., Rubinstein, M. and Patchornik, A. Selective covalent binding of methionyl containing peptides and proteins to water insoluble polymeric reagents and their regeneration. *Biochemistry* **16**, 1424, 1977.
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11. Rubinstein, M., Stein, S. and Udenfriend, S. Isolation and characterization of opioid peptides from rat pituitary: b-Endorphin. *Proc. Natl. Acad. Sci. USA* **74**, 4969, 1977.
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13. Rubinstein, M., Stein, S. and Udenfriend, S. Characterization of proopiocortin, a precursor to opioid peptides and corticotropin. *Proc. Natl. Acad. Sci. USA* **75**, 669, 1978.

14. Stein, S., Rubinstein, M., and Udenfriend, S. Ultramicro analysis of peptides. *Psychopharmacology Bulletin (NIMH)* **14**, 29, 1978.
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26. Evinger, M., Rubinstein, M. and Pestka, S. Antiproliferative and antiviral activities of human leukocyte interferons. *Archives of Biochemistry and Biophysics* **210**, 319-329, 1981.



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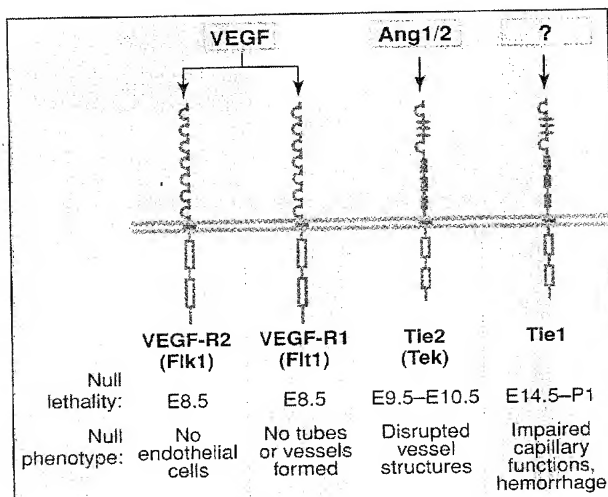
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# Signaling Vascular Morphogenesis and Maintenance

Douglas Hanahan

Blood vessels are constructed by two processes: vasculogenesis, whereby a primitive vascular network is established during embryogenesis from multipotential mesenchymal progenitors, and angiogenesis, in which preexisting vessels (both in embryo and adult) send out capillary sprouts to produce new vessels (1–3). Endothelial cells are centrally involved in each process: They migrate and proliferate and then assemble into tubes with tight cell-cell connections to contain the blood. Peri-endothelial support cells are recruited to encase the endothelial tubes, providing maintenance and modulatory functions to the vessels; such cells include pericytes for small capillaries, smooth muscle cells for larger vessels, and myocardial cells in the heart.

The establishment and remodeling of blood vessels is controlled by paracrine signals, many of which are protein ligands that bind and modulate the activity of transmembrane receptor tyrosine kinases (RTKs). This realization has emerged from the discovery and analysis of RTKs expressed on endothelial cells and of their ligands. Our basic view of RTK signaling has come from studies (performed largely in fibroblasts) of ligand-dependent autophosphorylation and activation of the branched Ras pathways. The results suggest that most RTKs are similarly coupled into the intracellular signal transduction cascade and are capable of inducing cell proliferation. However, the lessons from endothelial cells present a far more complicated picture. This complexity is highlighted by an article in this issue (4, page 55) of a ligand, called angiopoietin-2 (Ang2), that interferes with the kinase activity of an endothelial cell-selective RTK named Tie2. Remarkably, this inhibition of Tie2 kinase activity does not block endothelial cell pro-



**Fig. 1. Lessons from gene-knockout mice.** The endothelial cell-selective RTKs VEGF-R1, VEGF-R2, Tie1, and Tie2 have all been ablated in gene-knockout mice. Each RTK knockout produced embryonic lethality with vascular defects. However, their distinctive phenotypes indicate that each of these tyrosine kinases controls a specific, complementary function in endothelial cells that collectively can account for a significant part of endothelial cell morphogenesis into functional vessels. See (22) for receptor structures.

liferation and angiogenesis, but rather facilitates it.

Tie2 has two ligands, Ang1 and Ang2. Another major regulator of vasculogenesis and angiogenesis is vascular endothelial growth factor (VEGF, also called vascular permeability factor, VPF). VEGF signaling is itself mediated by two other endothelial cell-selective RTKs, called VEGF-R1 and VEGF-R2 (Flt1 and Flk1/KDR, respectively).

Mice carrying homozygous disruptions in the two highly homologous VEGF receptors die in mid-gestation of acute vascular defects, implicating each in vasculogenesis and developmental angiogenesis. However, the phenotypes are distinct—and instructive (see figure 1). VEGF-R2 knockout mice, which die by embryonic day 8.5 (E8.5), lack both endothelial cells and a developing hematopoietic system, implicating VEGF as a signal in the determination first of a hematopoietic progenitor and then of endothelial cells (5). In contrast, VEGF-R1 knockout mice, which also die around E8.5, have normal hematopoietic progenitors and

abundant endothelial cells, which migrate and proliferate but do not assemble into tubes and functional vessels (6). Thus, these highly homologous RTKs send distinctive signals in endothelial cells.

Tie2 knockout mice die somewhat later in embryogenesis (E9.5 to E10.5). The Tie2 null phenotype is distinct from that of the VEGF receptor knockouts and is also informative. Endothelial cells are present in normal numbers and are assembled into tubes, but the vessels are immature, lacking branching networks and proper organization into large and small vessels (7, 8). There is also an absence of the angiogenesis that vascularizes the neuroectoderm by capillary sprouting from the primitive vascular network (or plexus). Notably, the vessels that do form lack an intimate encapsulation by peri-endothelial support cells. In the heart, the endocardium and myocardium do not show tight association and structural complexity; rather, the endocardial cells have aberrant, rounded shapes, are only loosely attached to the surrounding basement membrane, and in many locations are disconnected from myocardial cells. Similar defects in vessel architecture are evident in other tissues. Thus, the Tie2 tyrosine kinase appears to control the capability of endothelial cells to recruit stromal cells to encase the endothelial tubes so as to stabilize the structure and modulate the function of blood vessels.

The fourth endothelial cell-selective RTK, Tie1, is remarkably similar in structure to Tie2 and appears to control another aspect of vessel integrity. Knockout mice lacking Tie1 die over a variable period, ranging from E14.5 to birth, of edema and hemorrhage, implicating the Tie1 signal in control of fluid exchange across capillaries and in hemodynamic stress resistance (8, 9).

New insights into the surprising concept that the Tie2 RTK is primarily coupled into a signal transduction circuit that elicits vessel maturation and maintains vessel integrity comes from functional analyses of the angiopoietins that bind to Tie2 and modulate its activity (4, 10, 11). The Tie2 ligands Ang1 and Ang2 are both ~75-kD secreted proteins with considerable sequence homology; each contains a coiled-coil and a fibrinogen-like domain. Both bind to the Tie2 receptor with similar affinity, and neither binds to the related receptor Tie1. Yet the effects on Tie2 are distinctive, as are the expression patterns in the mouse. Ang1 induces autophosphorylation of Tie2 in cultured endothelial cells. In marked contrast, Ang2, which binds with similar affinity, does not induce receptor phosphorylation. Rather, it can competitively inhibit Ang1-induced kinase activation of the Tie2 receptor. The

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Ang2 presents a negative signal to Tie2, a remarkable observation given its high homology to Ang1.

Moreover, this distinctive effect is apparently endothelial cell-specific. If a modified Tie2 is forcibly expressed in 3T3 fibroblasts, both Ang1 and Ang2 induce receptor phosphorylation and yet do not stimulate fibroblast proliferation. Similarly, Ang1-induced autophosphorylation of Tie2 does not affect endothelial cell growth in culture, consistent with the Tie2 knockout phenotype, which indicated that Tie2 is not required for endothelial cell proliferation during vasculogenesis. Functional studies in transgenic and gene-knockout mice support the notion that Ang1 signals Tie2 to recruit support cells, and that Ang2 inhibits this capability. Gene-knockout mice that lack Ang1 die with similar vascular defects to the Tie2 knockout mice (11). Transgenic mice overexpressing the negative ligand Ang2 in endothelial cells also die during embryogenesis, again with similar vascular defects (4). Thus, overexpression of Ang2 phenocopies loss of Ang1 expression, consistent with their opposite activities. Collectively, the data argue that Ang1 is the major physiological ligand for Tie2's functional role in recruiting and sustaining peri-endothelial support cells, whereas Ang2 serves to block this function, thereby relaxing these intimate and critical associations.

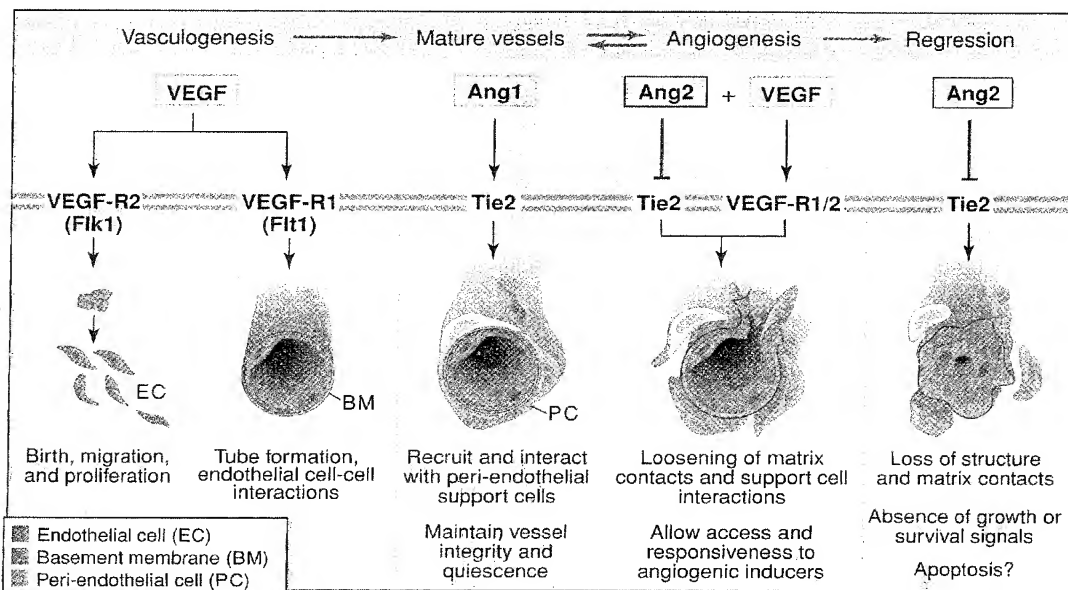
Why then does Ang2 exist? No doubt the Ang2 gene-knockout phenotype will prove illuminating. Meanwhile, the prevailing evidence suggests that Ang2 allows vascular remodeling, and in particular angiogenesis, processes that may be restricted by the encapsulation with basement membrane and peri-endothelial support cells. The clues come from the expression patterns of Ang1 and Ang2.

Ang1 is widely expressed both in the embryo and in the adult (4, 10). Ang2 is also widely expressed in the embryo. However, its expression pattern in the adult (in a limited survey) is provocative: Ang2 is selectively expressed in ovary, uterus, and placenta, the three tissues subject to physiological angiogenesis (4). The possible association of Ang2 with adult angiogenesis was investigated during ovulation, which is marked by distinctive phases of vascular quiescence, angiogenesis, and vascular regression. Ang2 expression in

these stages was compared to that of Ang1 and VEGF. In early follicles, the vasculature is quiescent, and Ang1 is expressed, with little or no VEGF or Ang2 expression. In late pre-ovulatory follicles and in the postovulatory corpus luteum, where angiogenesis is ongoing, both VEGF and Ang2 are up-regulated, while Ang1 expression persists. Notably, Ang2 expression appears punctate, or focal. Finally, in nonproductive follicles, which show vascular regression, Ang2 is expressed at uniformly high levels, provocatively in the absence of VEGF expression.

These patterns of expression, when considered in the context of the lessons from the various gene-knockout phenotypes (see Fig. 1), collectively suggest a model for control of vasculogenesis, vessel maturation and maintenance, angiogenesis, and regression, as illustrated in Fig. 2. The Ang1/Tie2 circuit appears to mediate vessel maturation from simple endothelial tubes into more elaborate vascular structures composed of several cell types, and the maintenance of those vessels via the cell-cell and cell-matrix associations

that produce them. Consequently, Ang1 may also help preserve endothelial cell quiescence. Focal expression of Ang2 evidently blocks the Ang1/Tie2 signal, resulting in a loosening of this tight vascular structure and thereby exposing the endothelial cells to activating signals from angiogenesis inducers, including VEGF. If VEGF (or another angiogenesis inducer) is present, the endothelial cells become activated to migrate and proliferate, producing new capillary sprouts and in turn tubes. One can envision that the more uniform presence of Ang1 allows a shift in the local balance of Ang1/Ang2 back in favor of Ang1, to effect maturation and stabilization of the newly formed vessels. Thus, there appears to be a collaboration between VEGF, Ang2, and Ang1 to elicit angiogenesis. In contrast, vascular regression is associated with very high-level expression of Ang2 in the absence of the activating (survival) signal from VEGF, presumably overwhelming the Ang1 signal and thereby producing catastrophic detachment from matrix and support cells, most likely with consequent apoptosis.



**Fig. 2. Regulation of vascular morphogenesis, maintenance, and remodeling by RTKs and their ligands.** A model for regulation of the vascular endothelium manifested by the prototypical angiogenesis factor VEGF and a new class of angiogenic regulators, Ang1 and Ang2. All three ligands bind to RTKs that have similar cytoplasmic signaling domains. Yet their downstream signals elicit distinctive cellular responses. Only VEGF binding to the VEGF-R2 sends a classical proliferative signal. When first activated in embryogenesis, this interaction induces the birth and proliferation of endothelial cells. In contrast, VEGF binding to VEGF-R1 elicits endothelial cell-cell interactions and capillary tube formation, a process that follows closely proliferation and migration of endothelial cells. Ang1 binding to the Tie2 RTK recruits and likely maintains association of peri-endothelial support cells (pericytes, smooth muscle cells, myocardiocytes), thus solidifying and stabilizing a newly formed blood vessel. The newly discovered Ang2, although highly homologous to Ang1, does not activate the Tie2 RTK; rather, it binds and blocks kinase activation in endothelial cells. The Ang2 negative signal causes vessel structures to become loosened, reducing endothelial cell contacts with matrix and disassociating peri-endothelial support cells. This loosening appears to render the endothelial cells more accessible and responsive toward the angiogenic inducer VEGF (and likely to other inducers). Finally, Ang2 is expressed at uniformly high levels in vascular regression in nonproductive ovarian follicles; the lack of VEGF coexpression suggests that loosening of cell-matrix interactions in the absence of a growth or survival signal elicits endothelial cell death, likely by apoptosis. No doubt additional factors play into these distinctive states, including the emerging class of angiogenesis inhibitors that directly block endothelial cell proliferation and migration.



The work by Maisonpierre *et al.* (4) introduces an inhibitory ligand, Ang2, for the Tie2 regulatory RTK that normally helps maintain vascular integrity. A model is emerging of a regulatory mechanism through which blood vessels are constructed, maintained, remodeled, and eliminated. Other factors are also involved, during embryonic vasculogenesis and angiogenesis, and for physiological and pathological angiogenesis in the adult. For example, gene-knockout mice have also implicated a G protein-coupled receptor in vascular regulation, because in the absence of  $G_{\alpha 13}$ , embryos die at E8.5 to E9.5, with defects in vascular assembly and angiogenesis (12). Moreover, a reciprocal paracrine signal has been revealed by gene-knockout mice that lack the genes for neuregulin, which is expressed in the endocardium of the heart, and the ErbB-2/3/4 RTKs, which are expressed in myocardium. Knockouts of this ligand or of its receptors produces defects in the developing heart analogous to those observed when Ang1 or Tie2 is missing (13–16). A similar role is suggested by gene-knockout mice for platelet-derived growth factor and its receptors in other tissue vasculature (17, 18). Furthermore, there is clear evidence in the adult for additional angiogenesis inducers and for an increasing number of angiogenesis inhibitors that act directly on endothelial cells (19–21). Thus, regulation of angiogenesis in ovulation and implantation, in wound healing, and in chronic pathological situations such as tumor progression will indeed be complex, but tractable using the power of animal models.

That complexity notwithstanding, the evidence is compelling that VEGF and the angiopoietins, and their cognate receptors, are critical components of the vascular regulatory machinery. It will be of particular interest to establish the possible contributions of Ang2 to tumor angiogenesis, whereby the quiescent vasculature (likely maintained in part by Ang1/Tie2) is activated to elicit and chronically effect the angiogenic phenotype that accompanies tumor growth and metastasis. Ang2 could well serve as an initial angiogenic signal, locally opening up the vessel structure to allow protease degradation of the basement membrane surrounding the endothelium and accessibility to that endothelium by angiogenesis inducers such as VEGF, thereby eliciting capillary sprouting and in turn new blood vessels that sustain a tumor as it expands.

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#### CONDENSED MATTER PHYSICS

## Low-Energy Excitations in High-Temperature Superconductors

Patrick A. Lee

Many common metals undergo a phase transition at some low critical temperature  $T_c$  and become superconductors. This phenomenon was explained in the famed BCS (Bardeen-Cooper-Schrieffer) theory (1) in terms of the coupling of electrons into Cooper pairs. In these materials, the superconducting ground state gains binding energy as a result of the pairing. It costs a finite amount of energy, called the energy gap, to break up the pairs to form excitations called quasiparticles. These are quantum mechanical superpositions of the electron and the hole excitations of the metallic state. These quasiparticles are interesting objects in their own right, but at low temperatures their number is exponentially small, and they hardly make their presence felt.

Until a decade ago, it was generally believed that the relative angular momentum of the electrons in Cooper pairs is *s* wave. Consequently, the energy gap is independent of the momentum *k* of the quasiparticle. This independence is in contrast to the only other case of Cooper pairing between fermions known at the time, namely, the superfluid state in liquid  $^3\text{He}$  (2). There the pairing state is anisotropic (*p* wave), and the energy gap may vanish along some momentum direction. These nodal points in the energy gap play an important role because quasiparticles near the nodes can be created in large numbers and they are responsible for the low-temperature properties. Today the prevailing view is that in high- $T_c$  superconductors, the Cooper pairs are formed in the *d*-

wave state. The role of the quasiparticles is then more analogous to that in superfluid  $^3\text{He}$ , and experimentalists are beginning to study them in detail. On page 83 of this issue, Krishana *et al.* (3) report a particularly interesting example of this class of experiments.

Although the *s*-wave BCS pairing theory works perfectly for practically all metallic superconductors, even a decade ago there was suspicion that a few superconductors are exceptions. The main evidence was that the low-temperature properties are not exponentially activated, pointing to the existence of nodes in the energy gap. The suspected unconventional superconductors were the so called "heavy fermion" superconductors (4) containing rare-earth ions such as Ce or U and a number of organic superconductors (5). The field of unconventional superconductivity took a dramatic turn in 1986 with the discovery of high-temperature superconductivity in cuprates (6). Apart from the high  $T_c$ , the metallic state (and to a lesser extent, the superconducting state) of these materials is so unusual that it was widely speculated that the pairing state is not the conventional *s* wave. After a number of brilliant experiments (7), it is now widely accepted that the Cooper pairs in the cuprates are formed in the *d*-wave symmetry. This symmetry implies that the energy gap vanishes at four points on the Fermi surface of these two-dimensional materials. From

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